

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING

ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/523,400A
Source: J.FWP
Date Processed by STIC: 12/06/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<**<http://www.uspto.gov/ebc/efs/downloads/documents.htm>**> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	SERIAL NUMBER: <u>10/523,400A</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 _____ Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 _____ Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 _____ Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters , instead.	
4 _____ Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 _____ Variable Length	Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 _____ PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 _____ Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 _____ Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional , please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 _____ Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 _____ Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or <u>scientific name (Genus/species)</u> . <220>-<223> section is required when <213> response is Unknown or Artificial Sequence. (see item 11 below)	
11 _____ Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of Sequence Rules	
12 _____ PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 _____ Misuse of n/Xaa	"n" can only represent a single <u>nucleotide</u> ; "Xaa" can only represent a single <u>amino acid</u>	



IFWP

RAW SEQUENCE LISTING

DATE: 12/06/2006

PATENT APPLICATION: US/10/523,400A

TIME: 09:38:18

Input Set : A:\US10523400-seq list(rev12-01-06).txt

Output Set: N:\CRF4\12062006\J523400A.raw

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5 <110> APPLICANT: Bernard Pau
7 <120> TITLE OF INVENTION: Specific antibodies for diagnosing heart failure
9 <130> FILE REFERENCE: P70365US0
11 <140> CURRENT APPLICATION NUMBER: US 10/523,400A
12 <141> CURRENT FILING DATE: 2005-02-03
14 <150> PRIOR APPLICATION NUMBER: PCT/FR03/02483
15 <151> PRIOR FILING DATE: 2003-08-07
17 <150> PRIOR APPLICATION NUMBER: FR 0210063
18 <151> PRIOR FILING DATE: 2002-08-07
20 <160> NUMBER OF SEQ ID NOS: 124
22 <170> SOFTWARE: PatentIn version 3.1
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 108
26 <212> TYPE: PRT
27 <213> ORGANISM: Homo sapiens : proBNP(1-108)
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32 His Pro Leu Gly Ser Pro Gly Ser Ala Ser Asp Leu Glu Thr Ser Gly
33 1 5 10 15
36 Leu Gln Glu Gln Arg Asn His Leu Gln Gly Lys Leu Ser Glu Leu Gln
37 20 25 30
40 Val Glu Gln Thr Ser Leu Glu Pro Leu Gln Glu Ser Pro Arg Pro Thr
41 35 40 45
44 Gly Val Trp Lys Ser Arg Glu Val Ala Thr Glu Gly Ile Arg Gly His
45 50 55 60
48 Arg Lys Met Val Leu Tyr Thr Leu Arg Ala Pro Arg Ser Pro Lys Met
49 65 70 75 80
52 Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp Arg Ile Ser Ser
53 85 90 95
56 Ser Ser Gly Leu Gly Cys Lys Val Leu Arg Arg His
57 100 105
60 <210> SEQ ID NO: 2
61 <211> LENGTH: 32
62 <212> TYPE: PRT
63 <213> ORGANISM: Homo sapiens : proBNP(77-108)
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69 Ser Pro Lys Met Val Gln Gly Ser Gly Cys Phe Gly Arg Lys Met Asp
70 1 5 10 15
73 Arg Ile Ser Ser Ser Ser Gly Leu Gly Cys Lys Val Leu Arg Arg His
74 20 25 30
77 <210> SEQ ID NO: 3
78 <211> LENGTH: 76
79 <212> TYPE: PRT
80 <213> ORGANISM: Homo sapiens : proBNP(1-76)

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RAW SEQUENCE LISTING

DATE: 12/06/2006

PATENT APPLICATION: US/10/523,400A

TIME: 09:38:18

Input Set : A:\US10523400-seq list(rev12-01-06).txt

Output Set: N:\CRF4\12062006\J523400A.raw

83 <400> SEQUENCE: 3

85 His Pro Leu Gly Ser Pro Gly Ser Ala Ser Asp Leu Glu Thr Ser Gly

86 1 5 10 15

88 Leu Gln Glu Gln Arg Asn His Leu Gln Gly Lys Leu Ser Glu Leu Gln

89 20 25 30

91 Val Glu Gln Thr Ser Leu Glu Pro Leu Gln Glu Ser Pro Arg Pro Thr

92 35 40 45

94 Gly Val Trp Lys Ser Arg Glu Val Ala Thr Glu Gly Ile Arg Gly His

95 50 55 60

97 Arg Lys Met Val Leu Tyr Thr Leu Arg Ala Pro Arg

98 65 70 75

101 <210> SEQ ID NO: 4

102 <211> LENGTH: 16

103 <212> TYPE: PRT

104 <213> ORGANISM: Artificial Sequence

107 <220> FEATURE:

108 <221> NAME/KEY: MOD_RES

109 <222> LOCATION: (1)..(1)

110 <223> OTHER INFORMATION: chemically synthesized

112 <400> SEQUENCE: 4

114 Tyr Thr Leu Arg Ala Pro Arg Ser Pro Lys Met Val Gln Gly Ser Gly

115 1 5 10 15

118 <210> SEQ ID NO: 5

119 <211> LENGTH: 6

120 <212> TYPE: PRT

121 <213> ORGANISM: Artificial Sequence

124 <220> FEATURE:

125 <221> NAME/KEY: MOD_RES

126 <222> LOCATION: (1)..(1)

127 <223> OTHER INFORMATION: chemically synthesized

129 <400> SEQUENCE: 5

131 Arg Ala Pro Arg Ser Pro

132 1 5

135 <210> SEQ ID NO: 6

136 <211> LENGTH: 8

137 <212> TYPE: PRT

138 <213> ORGANISM: Artificial Sequence

141 <220> FEATURE:

142 <221> NAME/KEY: MOD_RES

143 <222> LOCATION: (1)..(1)

144 <223> OTHER INFORMATION: chemically synthesized

147 <400> SEQUENCE: 6

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150 1 5

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154 <211> LENGTH: 8

155 <212> TYPE: PRT

156 <213> ORGANISM: Artificial Sequence

159 <220> FEATURE:

RAW SEQUENCE LISTING

DATE: 12/06/2006

PATENT APPLICATION: US/10/523,400A

TIME: 09:38:18

Input Set : A:\US10523400-seq list(rev12-01-06).txt

Output Set: N:\CRF4\12062006\J523400A.raw

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168 1      5
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172 <211> LENGTH: 9
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174 <213> ORGANISM: Artificial Sequence
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179 <222> LOCATION: (1)..(1)
180 <223> OTHER INFORMATION: chemically synthesized
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186 1      5
189 <210> SEQ ID NO: 9
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191 <212> TYPE: PRT
192 <213> ORGANISM: Artificial Sequence
195 <220> FEATURE:
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197 <222> LOCATION: (1)..(1)
198 <223> OTHER INFORMATION: chemically synthesized
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203 Cys Gly Arg Ala Pro Arg Ser Pro Lys
204 1      5
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209 <212> TYPE: PRT
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215 <222> LOCATION: (1)..(1)
216 <223> OTHER INFORMATION: chemically synthesized
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221 Cys Gly Arg Ala Pro Arg Ser Pro Lys Met Val
222 1      5      10
225 <210> SEQ ID NO: 11
226 <211> LENGTH: 15
227 <212> TYPE: PRT
228 <213> ORGANISM: Artificial Sequence
231 <220> FEATURE:
232 <221> NAME/KEY: MOD_RES
233 <222> LOCATION: (1)..(1)
234 <223> OTHER INFORMATION: chemically synthesized
237 <400> SEQUENCE: 11
239 Cys Gly Arg Ala Pro Arg Ser Pro Lys Met Val Gln Gly Ser Gly

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RAW SEQUENCE LISTING

DATE: 12/06/2006

PATENT APPLICATION: US/10/523,400A

TIME: 09:38:18

Input Set : A:\US10523400-seq list(rev12-01-06).txt

Output Set: N:\CRF4\12062006\J523400A.raw

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255 <223> OTHER INFORMATION: chemically synthesized
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261 Arg Ala Pro Arg Ser Pro Gly Cys
262 1          5
265 <210> SEQ ID NO: 13
267 <211> LENGTH: 8
269 <212> TYPE: PRT
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275 <220> FEATURE:
277 <221> NAME/KEY: MOD_RES
279 <222> LOCATION: (1)..(1)
281 <223> OTHER INFORMATION: chemically synthesized
285 <400> SEQUENCE: 13
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288 1          5
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303 <221> NAME/KEY: MOD_RES
305 <222> LOCATION: (1)..(1)
307 <223> OTHER INFORMATION: chemically synthesized
311 <400> SEQUENCE: 14
313 Cys Tyr Thr Leu Arg Ala Pro Arg Ser Pro Lys
314 1          5          10
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321 <212> TYPE: PRT
323 <213> ORGANISM: Artificial Sequence
326 <220> FEATURE:
327 <221> NAME/KEY: MOD_RES
328 <222> LOCATION: (1)..(1)
329 <223> OTHER INFORMATION: chemically synthesized
333 <400> SEQUENCE: 15
335 Cys His Arg Lys Met Val Leu Tyr Thr Leu Arg Ala Pro Arg Ser Pro
336 1          5          10          15
339 Lys
343 <210> SEQ ID NO: 16
345 <211> LENGTH: 17
347 <212> TYPE: PRT

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RAW SEQUENCE LISTING

DATE: 12/06/2006

PATENT APPLICATION: US/10/523,400A

TIME: 09:38:18

Input Set : A:\US10523400-seq list(rev12-01-06).txt

Output Set: N:\CRF4\12062006\J523400A.raw

349 <213> ORGANISM: Artificial Sequence
352 <220> FEATURE:
353 <221> NAME/KEY: MOD_RES
354 <222> LOCATION: (1)..(1)
355 <223> OTHER INFORMATION: chemically synthesized
359 <400> SEQUENCE: 16
361 Cys Tyr Thr Leu Arg Ala Pro Arg Ser Pro Lys Met Val Gln Gly Ser
362 1 5 10 15
365 Gly
369 <210> SEQ ID NO: 17
371 <211> LENGTH: 17
373 <212> TYPE: PRT
375 <213> ORGANISM: Artificial Sequence
378 <220> FEATURE:
379 <221> NAME/KEY: MOD_RES
380 <222> LOCATION: (1)..(1)
381 <223> OTHER INFORMATION: chemically synthesized
385 <400> SEQUENCE: 17
387 Cys Phe Thr Leu Arg Ala Pro Arg Ser Pro Lys Met Val Gln Gly Ser
388 1 5 10 15
391 Gly
395 <210> SEQ ID NO: 18
397 <211> LENGTH: 17
399 <212> TYPE: PRT
401 <213> ORGANISM: Artificial Sequence
404 <220> FEATURE:
405 <221> NAME/KEY: MOD_RES
406 <222> LOCATION: (1)..(1)
407 <223> OTHER INFORMATION: chemically synthesized
411 <400> SEQUENCE: 18
413 Cys Phe Ser Ile Arg Ala Pro Arg Ser Pro Lys Met Val Gln Gly Ser
414 1 5 10 15
417 Gly
421 <210> SEQ ID NO: 19
423 <211> LENGTH: 17
425 <212> TYPE: PRT
427 <213> ORGANISM: Artificial Sequence
431 <220> FEATURE:
433 <221> NAME/KEY: MOD_RES
435 <222> LOCATION: (17)..(17)
437 <223> OTHER INFORMATION: bAla
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443 Cys Tyr Thr Leu Arg Ala Pro Arg Ser Pro Lys Met Val Gln Gly Ser
444 1 5 10 15
447 Ala
451 <210> SEQ ID NO: 20
453 <211> LENGTH: 17
455 <212> TYPE: PRT
457 <213> ORGANISM: Artificial Sequence

→ If 2137 Responses are Artificial or Unknown, pls explain the Source of genetic material. See Item 11 on Error Summary sheet.

VERIFICATION SUMMARY

DATE: 12/06/2006

PATENT APPLICATION: US/10/523,400A

TIME: 09:38:19

Input Set : A:\US10523400-seq list(rev12-01-06).txt

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